In the Claims:

Please amend Claim 39 as follows:

- 39. (Once amended) An isolated polypeptide having at least 80% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127);
- (b) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127); or
- (d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide; or
- the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263,

 wherein said polypeptide is capable of inhibiting neoplastic cell growth.
- 40. (Once amended) The isolated polypeptide of Claim 39 having at least 85% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127);
- (b) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide;
- (e)(c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127); or
- (d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide; or
- the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263, wherein said polypeptide is capable of inhibiting neoplastic cell growth.



- 41. (Once amended) The isolated polypeptide of Claim 39 having at least 90% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127);
- (b) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127); or
- (d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide; or
- the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263, wherein said polypeptide is capable of inhibiting neoplastic cell growth.
- 42. (Once amended) The isolated polypeptide of Claim 39 having at least 95% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127);
- (b) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127); or
- (d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide; or
- the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263, wherein said polypeptide is capable of inhibiting neoplastic cell growth.

- 43. (Once amended) The isolated polypeptide of Claim 39 having at least 99% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127);
- (b) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide,
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEO ID NO:127); or
- (d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide; or
- the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263, wherein said polypeptide is capable of inhibiting neoplastic cell growth.
- 44. (Once amended) An isolated polypeptide comprising:
- (a) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127);
- (b) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127); or
- (d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide; or
- the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263.
- 45. (Previously added) The isolated polypeptide of Claim 44 comprising the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127).



- 46. (Previously added) The isolated polypeptide of Claim 44 comprising the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide.
- 47. (Previously added) The isolated polypeptide of Claim 44 comprising the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127).
- 48. Cancel.
- 49. (Previously added) The isolated polypeptide of Claim 44 comprising the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263.
- (Previously added) A chimeric polypeptide comprising a polypeptide according to Claim 39 fused to a heterologous polypeptide.
 - 51. (Previously added) The chimeric polypeptide of Claim 50, wherein said heterologous polypeptide is an epitope tag or an Fc region of an immunoglobulin.